CRF<u>T</u> 1646





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PATENT APPLICATION: US/09/711,724A

DATE: 01/17/2003 TIME: 07:55:15

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PATENT APPLICATION: US/09/711,724A TIME: 07:55:15

Input Set: N:\Crf3\dbback2\Datahold\EFS\09711724\HMSU-P14-

DATE: 01/17/2003

006SubstituteSequence.txt

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    267 <212> TYPE: DNA
    268 <213> ORGANISM: Homo sapiens
    270 <220> FEATURE:
    271 <221> NAME/KEY: misc_feature
    272 <222> LOCATION: (1387)..(1389)
    273 <223> OTHER INFORMATION: n=a, c, g, or t
    275 <400> SEOUENCE: 6
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    278 ggactggcgt gcggaccggg cagggggttc gggaagagga ggcaccccaa aaagctgacc
    280 cctttagcct acaaqcaqtt tatccccaat qtqqccqaqa aqaccctagg cgccagcgga
                                                                               180
                                                                               240
    282 aggtatgaag ggaagatete cagaaactee gagegattta aggaacteae eeccaattae
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    284 aaccccgaca tcatatttaa ggatgaagaa aacaccggag cggacaggct gatgactcag
                                                                               360
    286 aggtgtaagg acaagttgaa cgctttggcc atctcggtga tgaaccagtg gccaggagtg
    288 aaactgcggg tgaccgaggg ctgggacgaa gatggccacc actcagagga gtctctgcac
                                                                               420
                                                                               480
    290 tacgagggcc gcgcagtgga catcaccacg tctgaccgcg accgcagcaa gtacggcatg
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    292 ctggcccgcc tggcggtgga ggccggcttc gactgggtgt actacgagtc caaggcacat
                                                                               600
    294 atccactgct cggtgaaagc agagaactcg gtggcggcca aatcgggagg ctgcttcccg
    296 ggctcggcca cggtgcacct ggagcagggc ggcaccaagc tggtgaagga cctgagcccc
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    298 ggggaccgcg tgctggcggc ggacgaccag ggccggctgc tctacagcga cttcctcact
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     300 ttcctggacc gcgacgacgg cgccaagaag gtcttctacg tgatcgagac gcgggagccg
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    302 egegagegee tgetgeteae egeegegeae etgetetttg tggegeegea caaegaeteg
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     304 gecacegggg agecegagge gteeteggge teggggeege etteeggggg egeaetgggg
     306 cctcgggcgc tgttcgccag ccgcgtgcgc ccgggccagc gcgtgtacgt ggtggccgag
                                                                               960
                                                                              1020
     308 cgtgacgggg accgccggct cctgcccgcc gctgtgcaca gcgtgaccct aagcgaggag
                                                                              1080
     310 geogegggeg cetaegegee geteaeggee eagggeacea tteteateaa eegggtgetg
                                                                              1140
    312 geotegtget acgeggteat egaggageae agetgggege acegggeett egegeeette
    314 cgcctggcgc acgcgctcct ggctgcactg gcgcccgcgc gcacggaccg cggcggggac
                                                                              1200
                                                                              1260
    316 ageggeggeg gggacegegg gggeggegge ggeagagtag cectaacege tecaggtget
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     318 geogaegete egggtgeggg ggeeaeegeg ggeateeaet ggtaetegea getgetetae
                                                                              1380
     320 caaataggca cetggeteet ggacagegag gecetgeace egetgggeat ggeggteaag
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     325 <211> LENGTH: 939
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    332 aaccagtggc ccggtgtgaa gctgcgggtg accgagggct gggacgagga cggccaccac
                                                                               120
                                                                               180
    334 teagaggagt ceetgeatta tgagggeege geggtggaea teaceacate agacegegae
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     336 cgcaataagt atggactgct ggcgcgcttg gcagtggagg ccggctttga ctgggtgtat
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     338 tacgagtcaa aggcccacgt gcattgctcc gtcaagtccg agcactcggc cgcagccaag
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    340 acggccgct gettecetge eggageceag gtacgcetgg agagtgggge gegtgtggee
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    342 ttgtcagccg tgaggccggg agaccgtgtg ctggccatgg gggaggatgg gagccccacc
                                                                               480
    344 ttcagcgatg tgctcatttt cctggaccgc gagccccaca ggctgagagc cttccaggtc
                                                                               540
    346 atcgagactc aggacccccc acgccgcctg gcactcacac ccgctcacct gctctttacg
    348 gctgacaatc acacggagcc ggcagcccgc ttccgggcca catttgccag ccacgtgcag
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    350 cctggccagt acgtgctggt ggctggggtg ccaggcctgc agcctgcccg cgtggcagct
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DATE: 01/17/2003 TIME: 07:55:15 PATENT APPLICATION: US/09/711,724A

Input Set : N:\Crf3\dbback2\Datahold\EFS\09711724\HMSU-P14-

006SubstituteSequence.txt

354 356 358 360 362 363	352 gtctctacac acgtggcct cggggcctac gcccgctca caaagcatgg gacactggtg 354 gtggaggatg tggtggcatc ctgcttcgcg gccgtggctg accaccacct ggctcagttg 356 gccttctggc ccctgagact ctttcacagc ttggcatggg gcagctggac cccgggggag 358 ggtgtgcatt ggtaccccca gctgctctac cgcctgggc gtctcctgct agaagagggc 360 agcttccacc cactgggcat gtccggggca gggagctga 362 <210> SEQ ID NO: 8 363 <211> LENGTH: 425 364 <212> TYPE: PRT										720 780 840 900 939						
			RGANI EQUEN			lus g	gallu	ıs									
	Met					Leu	Leu	Thr	Arg	Ile 10	Leu	Leu	Val	Gly	Phe 15	Ile	
		Ala	Leu	Leu 20	-	Ser	Ser	Gly	Leu 25		Cys	Gly	Pro	Gly 30	Arg	Gly	
	Ile	Gly	Lys 35		Arg	His	Pro	Lys 40		Leu	Thr	Pro	Leu 45		Tyr	Lys	
	Gln	Phe 50		Pro	Asn	Val	Ala 55		Lys	Thr	Leu	Gly 60	-	Ser	Gly	Arg	
			Gly	Lys	Ile	Thr 70		Asn	Ser	Glu	Arg 75		Lys	Glu	Leu	Thr 80	
		Asn	Tyr	Asn	Pro 85		Ile	Ile	Phe	Lys 90		Glu	Glu	Asn	Thr 95	Gly	
	Ala	Asp	Arg	Leu 100		Thr	Gln	Arg	Cys 105	Lys	Asp	Lys	Leu	Asn 110	Ala	Leu	
	Ala	Ile	Ser 115		Met	Asn	Gln	Trp 120	Pro	Gly	Val	Lys	Leu 125	Arg	Val	Thr	
	Glu	Gly 130	Trp	Asp	Glu	Asp	Gly 135	His	His	Ser	Glu	Ġlu 140	Ser	Leu	His	Tyr	
	Glu 145	Gly	Arg	Ala	Val	Asp 150	Ile	Thr	Thr	Ser	Asp 155	Arg	Asp	Arg	Ser	Lys 160	
409 410	Tyr	Gly	Met	Leu	Ala 165	Arg	Leu	Ala	Val	Glu 170	Ala	Gly	Phe	Asp	Trp 175	Val	
413 414	Tyr	Tyr	Glu	Ser 180	Lys	Ala	His	Ile	His 185	Cys	Ser	Val	Lys	Ala 190	Glu	Asn	
	Ser	Val	Ala 195	Ala	Lys	Ser	Gly	Gly 200	Cys	Phe	Pro	Gly	Ser 205	Ala	Thr	Val	
421 422	His	Leu 210	Glu	His	Gly	Gly	Thr 215	Lys	Leu	Val	Lys	Asp 220	Leu	Ser	Pro	Gly	
			Val												Ser	Asp 240	
															Phe 255	Tyr	
	Val	Ile	Glu	Thr 260		Gln	Pro	Arg	Ala 265	Arg	Leu	Leu	Leu	Thr 270	Ala	Ala	
	His	Leu	Leu 275		Val	Ala	Pro	Gln 280		Asn	Gln	Ser	Glu 285	Ala	Thr	Gly	
		Thr 290		Gly	Gln	Ala	Leu 295		Ala	Ser	Asn	Val 300		Pro	Gly	Gln	
			Tyr	Val	Leu	Gly		Gly	Gly	Gln	Gln		Leu	Pro	Ala	Ser	

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/711,724A

DATE: 01/17/2003 TIME: 07:55:16

Input Set : N:\Crf3\dbback2\Datahold\EFS\09711724\HMSU-P14-

006SubstituteSequence.txt

Output Set: N:\CRF4\01172003\I711724A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:6; N Pos. 1387,1388,1389
Seq#:18; N Pos. 15,32,36
Seq#:19; N Pos. 24,27
Seq#:20; N Pos. 13,16,19,23,27
Seq#:30; N Pos. 6,23,27
Seq#:31; N Pos. 4,7,10,14,19,22
Seq#:38; N Pos. 20,23
Seq#:39; N Pos. 11,26
Seq#:40; Xaa Pos. 7,9,44,85,93,98,112,132,137,139,181,183,185,186,189,191
Seq#:40; Xaa Pos. 196,200,206,207,209,211,212,216,217,219
Seq#:41; Xaa Pos. 7,8,9,12,13,14,17,19,22,27,29,30,31,33,40,41,44,45,46,48
Seq#:41; Xaa Pos. 53,54,71,79,83,84,85,87,95,100,107,114,115,116,125,134
Seq#:41; Xaa Pos. 135,139,141,157,158,160,162,166,167

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/711,724A

DATE: 01/17/2003 TIME: 07:55:16

Input Set : N:\Crf3\dbback2\Datahold\EFS\09711724\HMSU-P14-

006SubstituteSequence.txt

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L:322 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:1380
L:1190 M:283 W: Missing Blank Line separator, <220> field identifier
L:1200 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:1216 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
L:1232 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
L:1347 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0
L:1363 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0
L:1363 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:0
L:1617 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:0
L:1762 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:0
L:1768 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:32
L:1777 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:80
L:1780 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:96
L:1786 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:128
L:1795 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:176
L:1798 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:192
L:1801 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:208
L:2040 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:0
L:2043 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:16
L:2046 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:32
L:2049 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:48
L:2052 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:64
L:2055 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:80
L:2058 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:96
L:2061 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:112
L:2064 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:128
L:2067 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:144
L:2070 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:160
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